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Identification and Serotyping of *Microsporium canis* Isolates by Monoclonal Antibodies

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Hybridoma cells were produced by fusing mouse myeloma cells with spleen cells from mice primed with an exoantigen of *Microsporium canis*. Three clones produced antibodies which were examined by the Western blot technique for their potential usefulness in the identification of *M. canis* isolates and differentiation of strains within the species. Based on reactions with immunological determinants, all of the *M. canis* isolates tested presented either species- or strain-specific domains. Monoclonal antibodies proved to be useful reagents for the identification of *M. canis* isolates and for the differentiation of strains within the species. A purified antigen depleted of common antigenic determinants was obtained in affinity chromatography by using monoclonal antibody.

The application of hybridoma technology has proven to be useful in the identification of a wide range of microorganisms. Monoclonal antibodies may be used comparatively in different immunological assays for phylogenetic studies and species and strain identification.

In a previous report, we described the potential for serologically analyzing dermatophytes by precipitating monoclonal antibodies produced against an exoantigen of *Microsporium canis* (5).

This report deals with the properties of dermatophyte antigenic determinants as determined by the reaction of monoclonal antibodies with polypeptides separated by electrophoresis in polyacrylamide gels and then transferred electrically to nitrocellulose.

Purification of crude *M. canis* exoantigen by affinity chromatography with monoclonal antibodies might also be achieved to obtain antigens specific to *M. canis*.

MATERIALS AND METHODS

Cultures. Dermatophyte isolates were obtained from clinical specimens in our institute (Università Cattolica del Sacro Cuore; UCSC) or were graciously furnished by the Division of Mycotic Diseases, Centers for Disease Control, Atlanta, Ga. (see Tables 1 to 3). All the cultures were maintained in our collection at room temperature in sterile distilled water.

Reference antigens. *M. canis* CDC B2094 was the standard strain used for the production of the reference exoantigen and for mouse immunization. It was extracted in 24 h from a 7-day-old Sabouraud dextrose agar slant culture by using a merthiolate solution (1:5,000) in distilled water. The solution was filtered and concentrated 50-fold by lyophilization. The protein and carbohydrate contents of the reference antigen were adjusted to final concentrations of 2,500 and 550 µg/ml, respectively. The same procedure was used to produce exoantigens (protein content, 2,500 µg/ml) from all the cultures used in this study.

Immunization protocol. For the fusion, BALB/c mice were

immunized intraperitoneally with 0.1 ml of the reference antigen mixed with an equal amount of incomplete Freund adjuvant. The procedure was repeated once a week for 1 month. Three days before the fusion, the mice were reinoculated intraperitoneally with 0.1 ml of the reference antigen only. Before the fusion, blood samples were collected from each mouse and placed in microtiter plates sensitized with the *M. canis* reference antigen for enzyme-linked immunosorbent assay of antibody titer.

Production and screening of hybridomas. The procedures for fusion and selection of the hybridoma cells producing monoclonal antibodies against *M. canis* exoantigen were performed as described elsewhere (5).

Western blot analysis. Dermatophyte lyophilized exoantigens (5 ml) were solubilized with 0.5 ml of disruptor buffer consisting of 1 M Tris (pH 7), 60% (wt/vol) sucrose, 2% (wt/vol) sodium dodecyl sulfate, 5% (wt/vol) B-mercaptoethanol, and 0.02% bromphenol blue-saturated solution. A 50-µg sample of protein standards (150 µl), including RNase A (13,700 daltons), chymotrypsinogen (25,000 daltons), and ovalbumin (43,000 daltons), was added per ml of the antigen solution, and it was boiled for 5 min. Antigen samples were electrophoresed through 10% polyacrylamide gels (vertical gel system; Bethesda Research Laboratories, Inc., Gaithersburg, Md.). Electrophoresis was performed at 15 mA and continued for 18 h to allow the solvent front to reach the bottom of the gel. Immunoblotting of the gels was performed by a modification of the method of Towbin et al. (6). Briefly, the proteins were electrically transferred to nitrocellulose sheets (0.45-µm pore size; Schleicher & Schuell, Inc., Keene, N.H.) at 60 V for 3 h (Trans blot cell; Bio-Rad Laboratories, Richmond, Calif.). After 90 min, the transfer buffer (25 mM Tris base, 192 mM glycine, 20% methanol) was changed. The nitrocellulose sheets were cut into strips and incubated with 5% horse serum in phosphate-buffered saline (PBS; pH 7.6) for 30 min at room temperature. The strips were subsequently incubated in a shaker for 30 min at 37°C in a 1:10 dilution of hybridoma culture fluid. The strips were then washed with PBS, blocked once more with 5% horse serum in PBS, and incubated for 30 min in a shaker at 37°C with 3 ml of horseradish peroxidase-coupled

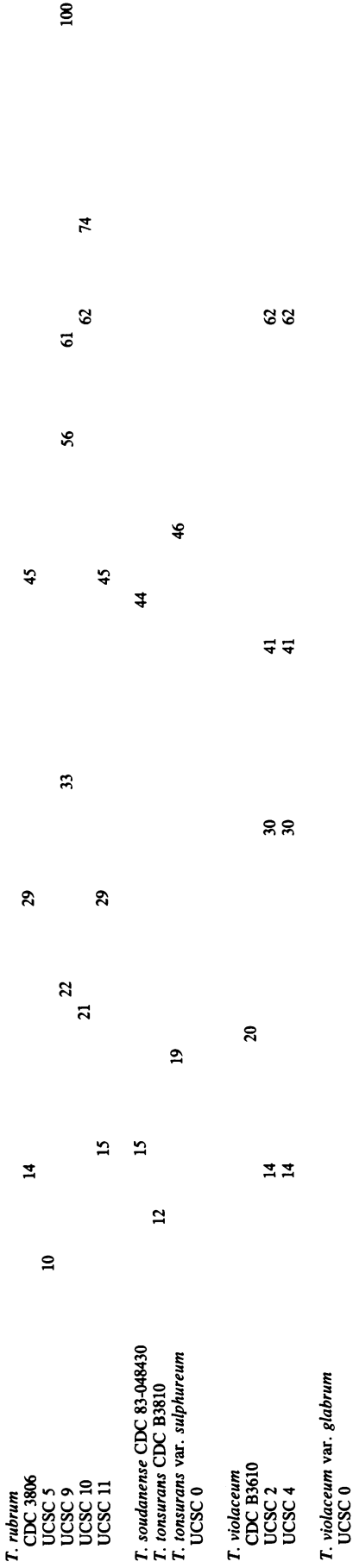
* Corresponding author.

TABLE 1. Western blot analysis of dermatophyte isolates with *M. canis* monoclonal antibody UCSC 3

Isolate	Molecular size (kDa)									
<i>Epidermophyton floccosum</i>										
CDC B3807										
UCSC 1	21	30	45	58	84					
UCSC 8	22	30	44	52	84					
<i>Microsporum audouinii</i>										
CDC B3800										
<i>M. canis</i>										
CDC B2094		28								
UCSC 1	19	21	23		74	74	80		95	
UCSC 2					74	74			95	
UCSC 4		25		60	74	74			95	
UCSC 6					74	74			95	
UCSC 8					74	74			95	
UCSC 10		30			74	74			95	
UCSC 12				48	74	74			95	
UCSC 14	19	25	37	45	74	74			95	
UCSC 52		25	40		74	74			95	
<i>M. cookei</i> CDC B3803										
<i>M. distortum</i> CDC B2174					72			89		
<i>M. equinum</i> CDC B2699										
<i>M. ferrugineum</i> CDC 83-056097					66					
<i>M. fulvum</i> UCSC 0		32	45		68		80			
<i>M. gallinae</i> CDC B3801										
UCSC 0		31	42	50	68			90	110	
<i>M. gypseum</i>										
CDC B3816	17									
UCSC 0	18	24				75	80			
UCSC 1		24	40			75				
UCSC 2		24	45	60	65			85		
UCSC 4		24	47							
<i>M. nanum</i> CDC B3815										
<i>M. persicolor</i> CDC B1923			38	49	68			98		
<i>M. racemosum</i> UCSC 0										
<i>M. ripariae</i> UCSC 0							78			
<i>Trichophyton megninii</i>										
UCSC 0	20	37					76	86		
<i>T. mentagrophytes</i> var. <i>erinacei</i>										
CDC B1865										
<i>T. mentagrophytes</i> var. <i>interdigitale</i>										
UCSC 3	15	22								
UCSC 5										
UCSC 6	15	22	42	62				92	120	
UCSC 7	14	24	46					92		
<i>T. mentagrophytes</i> var. <i>mentagrophytes</i>										
CDC B6209										
<i>T. mentagrophytes</i> var. <i>quinckeanum</i>	17	25	34	54	78					
CDC B2331			44	58						
			45							

TABLE 2. Western blot analysis of dermatophyte isolates with *M. canis* monoclonal antibody UCSC 7

Isolate	Molecular size (kDa)									
<i>Epidermophyton floccosum</i>										
CDC B3807	17									
UCSC 1									50	
UCSC 8									47	
<i>Microsporum audouinii</i>										
CDC B3800	14									
<i>M. canis</i>										
CDC B2094										
UCSC 1		22	25							98
UCSC 2									80	98
UCSC 4										98
UCSC 6										98
UCSC 8								64		98
UCSC 10										98
UCSC 12			27							98
UCSC 14										98
UCSC 52		24							75	98
<i>M. cookei</i> CDC B3803										
<i>M. distortum</i> CDC B2174	13							45		90
<i>M. equinum</i> CDC B2699										
<i>M. ferrugineum</i> CDC 83-056097	11	21								
<i>M. fulvum</i> UCSC 0	11		25							
<i>M. gallinae</i> CDC B3801										
UCSC 0		22						68		86
<i>M. gypseum</i>										
CDC B3816										
UCSC 0	11		25					40		
UCSC 1			25					34		
UCSC 2			25							
UCSC 4			25							
<i>M. nanum</i> CDC B3815										
<i>M. persicolor</i> CDC B1923										
<i>M. racemosum</i> UCSC 0										
<i>M. ripariae</i> UCSC 0		20								
<i>Trichophyton megninii</i>		21								
UCSC 0										
<i>T. mentagrophytes</i> var. <i>erinacei</i>										
CDC B1865										
<i>T. mentagrophytes</i> var. <i>interdigitale</i>										
UCSC 3	15									
UCSC 5										
UCSC 6	14								52	
UCSC 7	17									
<i>T. mentagrophytes</i> var. <i>mentagrophytes</i>										
CDC B6209	13									
<i>T. mentagrophytes</i> var. <i>quinckeanum</i>										
CDC B2331	17							37		84



changes, 200 ×, 4°C). The unbound, purified antigen was collected and lyophilized. The resin was used repeatedly without loss of binding activity.

Immunospot assay. A modification of the procedure of Hawkes et al. (2) for the assay of antibodies, based on the application of the antigen as a spot to nitrocellulose paper, was used to detect the reactivity of immunoadsorbed monoclonal antibody in affinity chromatography with either bound or unbound antigen.

RESULTS

Three hybridoma cell lines producing monoclonal antibodies of the immunoglobulin G class were selected according to their different reactivity with the reference antigen as detected by the Western blot technique. This procedure also permitted study of the properties of the dermatophyte proteins with the monoclonal antibodies. Our results were as follows.

M. canis monoclonal antibody UCSC 3 recognized all of the *M. canis* isolates tested by reaction with specific antigenic determinants of 74 and 95 kilodaltons (kDa) (Table 1). The detection of reactive antigenic determinants of different molecular weights in the different isolates within the species suggests the existence of numerous serotypes. No heterologous dermatophyte species, showing cross-reactivity, exhibited the simultaneous presence of the 74- and 95-kDa antigenic determinants.

M. canis monoclonal antibody UCSC 7 characterized single *M. canis* isolated by either single species-specific (98-kDa) or strain-specific antigenic determinants of different molecular weights (Table 2). It was interesting to note that the five *M. gypseum* isolates tested showed a common domain of 25 kDa.

M. canis monoclonal antibody UCSC 13 (Fig. 1) proved to be the most interesting and specific one. It reacted only with a single 16-kDa antigenic determinant in each *M. canis* isolate tested. This reaction was also shared by the *Microsporum distortum* isolate tested, which concurs with the identity of the two species as demonstrated previously by mating experiments (4). Cross-reactions, although at different molecular weights, were observed only in the *Microsporum ferrugineum*, *Microsporum gallinae*, and *Trichophyton soudanense* isolates tested. These species are considered strictly related morphologically (Table 3).

Affinity chromatography was successful in purifying one

TABLE 3. Western blot analysis of dermatophyte isolates^a with *M. canis* monoclonal antibody UCSC 13

Isolate	Molecular size (kDa)
<i>M. canis</i>	
CDC B2094	16
UCSC 1	16
UCSC 2	16
UCSC 4	16
UCSC 6	16
UCSC 8	16
UCSC 10	16
UCSC 12	16
UCSC 14	16
UCSC 52	16
<i>M. distortum</i> CDC B2174	16
<i>M. ferrugineum</i> CDC 83-056097	11
<i>M. gallinae</i> CDC B3801	23
<i>T. soudanense</i> CDC 83-048430	12

^a The other dermatophytes listed in Tables 1 and 2 were negative with this monoclonal antibody.

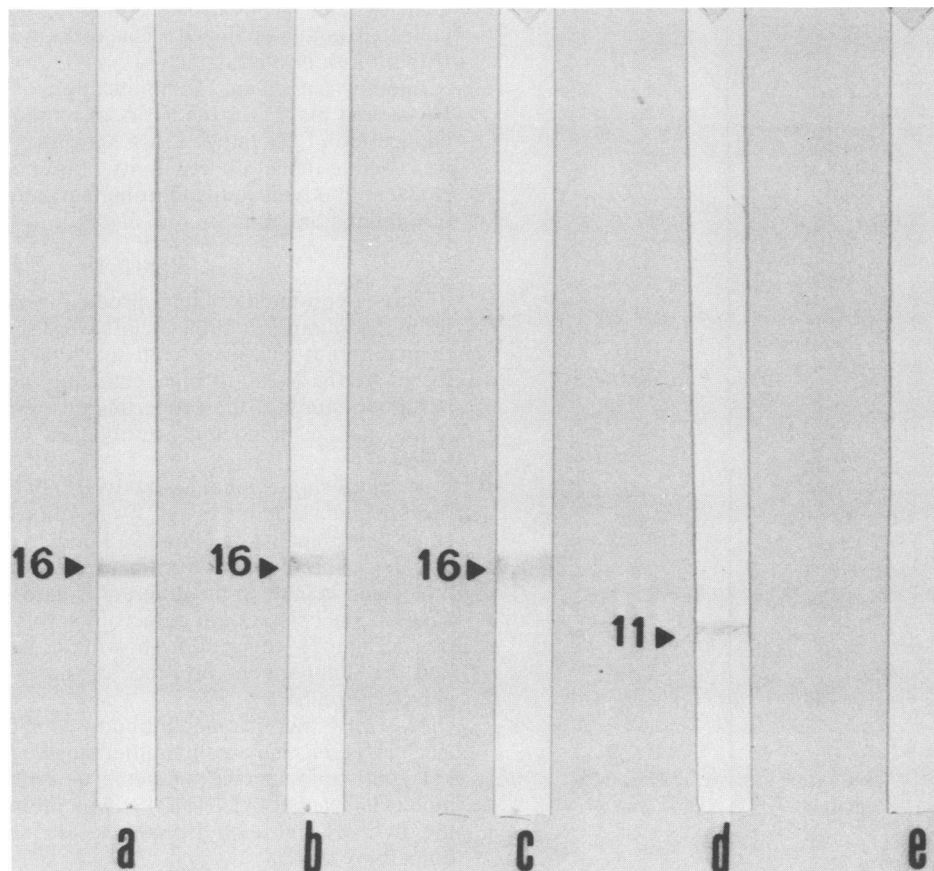


FIG. 1. Western blot analysis of dermatophyte isolates with *M. canis* monoclonal antibody UCSC 13 (molecular size expressed in kDa). Lanes: a, *M. canis* CDC B2094 exoantigen; b, *M. canis* UCSC 1 exoantigen; c, *M. canis* UCSC 12 exoantigen; d, *M. ferrugineum* CDC 83-056097 exoantigen; e, *Microsporum audouinii* CDC B3800 exoantigen.

component of the reference antigen. In a preliminary approach using *M. canis* monoclonal antibody UCSC 7 as an immunoabsorbent, it was possible to separate peak 1 (unadsorbed portion of the antigen) and peak 2 (adsorbed portion of the antigen) (Fig. 2).

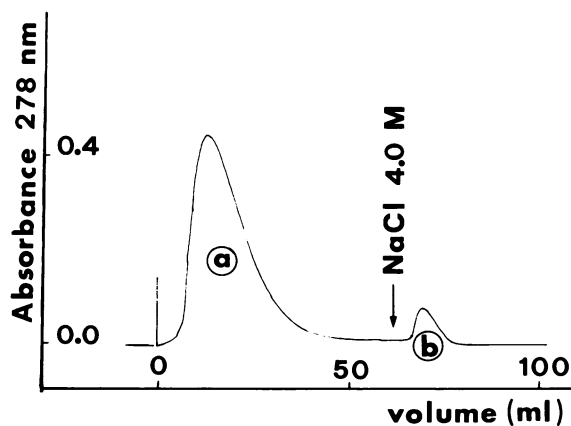


FIG. 2. Affinity chromatography of *M. canis* CDC B2094 total exoantigen against monoclonal antibody UCSC 7 coupled to Affigel 10 resin. Peak a represents the unadsorbed antigenic fraction, and peak b represents the fraction eluted by sodium chloride (4.0 M) in elution buffer (arrow).

The immunospot assay performed with the reference antigen in toto, peak 1 (unadsorbed antigen), and peak 2 (adsorbed antigen) with *M. canis* monoclonal antibody UCSC 7 showed the loss of reactivity of peak 1 (unadsorbed antigen) (Fig. 3).

Comparison by reverse-phase HPLC of the reference

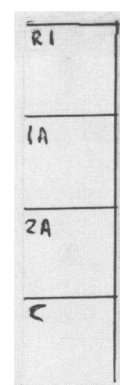


FIG. 3. Evaluation of reaction of in toto (R1), unadsorbed (1A), and adsorbed (2A) exoantigen of *M. canis* CDC B2094 by immunospot with *M. canis* monoclonal antibody UCSC 7. Antigenic fractions 1A and 2A were obtained after affinity chromatography of the exoantigen with *M. canis* monoclonal antibody UCSC 7. Control (C) was PBS used in place of the antigen.

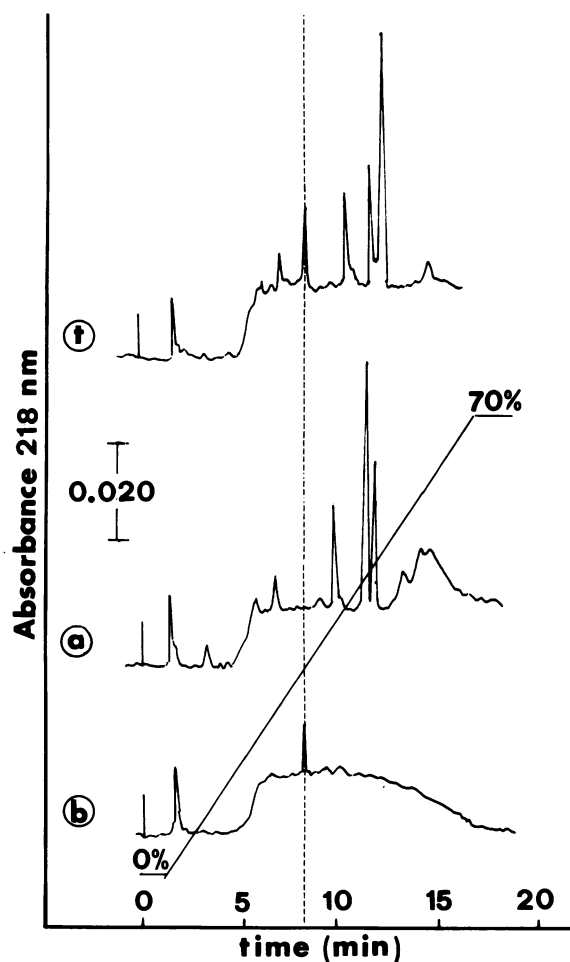


FIG. 4. Reverse-phase HPLC. t, *M. canis* CDC B2094 total exoantigen; a, unadsorbed antigenic fraction; b, the fraction eluted by sodium chloride (4.0 M) in elution buffer after affinity chromatography against monoclonal antibody UCSC 7 on Affigel 10 resin. The dashed line shows the position of the reactive antigenic determinant in the chromatographic pattern. The acetonitrile gradient applied in the chromatography is also indicated.

antigen and peak 1 (portion eluted) confirmed the disappearance of one peak of the original pattern (Fig. 4).

DISCUSSION

Our study clearly showed the enormous potential of monoclonal antibodies for the immunoidentification and typing of dermatophytes. The effectiveness of the monoclonal antibodies may be completely evaluated by using a highly

sensitive and analytical procedure, such as the Western blot technique. It is of primary importance, moreover, that the immunization antigen be an exoantigen which can very easily and rapidly be produced from the dermatophyte isolates to be identified.

When the dermatophyte antigens were electrophoretically separated in denaturing gels and then immobilized on nitrocellulose strips, we detected a greater diversity of monoclonal antibody reactivity to fungal proteins than when we used the technique of immunodiffusion of soluble nondenatured dermatophyte antigens (5). Analogous to the results obtained by Braun et al. (1) in the study of viral proteins, the preliminary advantage of the technique is in the detection of nonprecipitating antibody and of antibody to poorly soluble antigens.

The finding that multiple-molecular-weight components of the exoantigen reacted with monoclonal antibodies might suggest either modifications of a common protein or the polymerized states of a protein which was not separated under the denaturing conditions used in this study. Finally, the procedure of affinity chromatography by the immunoadsorption of nonspecific monoclonal antibodies might lead to the purification of antigens specific to *M. canis*.

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